

CERTIFICATE OF MAILING (37 CFR 1.8)

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

TECH CENTER 1600/2900

Art Unit : 1647
 Examiner : Sharon L. Turner, Ph.D.
 Applicant(s) : ME Gurney, AM Pauley, and J Li
 Serial Number : 09/328,877
 Filed : 9 June 1999
 For : Human Sel-10 Polypeptides and Polynucleotides that Encode Them

Commissioner of Patents and Trademarks
 Washington, DC 20231

TRANSMITTAL OF A RESPONSE TO A NON-FINAL ACTION (37 CFR 1.111)

Sir:

Transmitted herewith is a reply and/or amendment in the above-captioned application in response to the Examiner's action dated 14 June 2001.

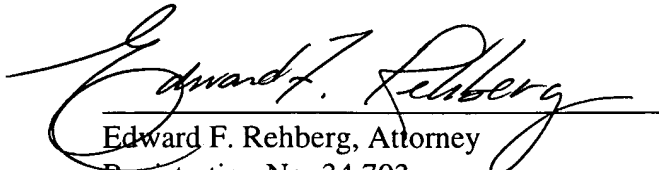
- ☒ [X] The reply and/or amendment is being filed under 37 CFR 1.8 and the required Certificate of Mailing appears above.
- ☐ [] An additional fee in the amount of \$ is required for the amended claims presented and has been calculated as shown in the attached sheet.

Please charge Deposit Account No. 21-0718 in the amount of the additional fee above, or such greater or lesser amount of excess fees for claims as the Commissioner determines is required by law. Triplicate copies of this sheet are enclosed.

EXTENSION OF TIME. In the event this paper is not filed prior to the time set for response, applicant(s) hereby petition for an extension of the period for filing the attached reply and/or amendment to the date of filing this paper, and hereby authorize the Commissioner to charge the extension fee as may be required by 37 CFR 1.17, to Deposit Account No. 21-0718. If for any

reason the extension requested above is insufficient to extend this period to the date of this paper, applicant(s) hereby petition for the revival of the above-captioned application as having been unintentionally abandoned and authorize the Commissioner to charge the required fees under 37 CFR 1.17 to Deposit Account No. 21-0718.

Respectfully submitted,


Edward F. Rehberg, Attorney
Registration No. 34,703

Date: 8-14-2001

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Enclosures:

Reply/Amendment

☐ Calculation of Additional Fees for Amended Claims



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Art Unit : 1647
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For : Human Sel-10 Polypeptides and Polynucleotides that Encode Them

Commissioner of Patents and Trademarks
Washington, DC 20231

Sir:

This is in response to the office action mailed 14 June 2001 in the above-identified application, the shortened statutory period for response being until 14 July 2001. A request for a one month extension of time for response is included herewith to extend the period for response to 14 August 2001.

Election

Group XIV is provisionally elected with traverse.

Background

The Examiner initially issued a four way restriction requirement asserting that claims directed to I. polynucleotides, II. polypeptides, III. antibodies and IV. methods of identifying agents were distinct. Applicant responded by electing group II with traverse. The Examiner responded by vacating the previously issued restriction requirement and imposed a 34 way restriction requirement.

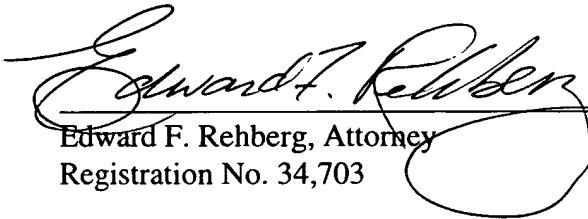
Traversal of the Restriction Requirement

The Applicant would point the Examiner to MPEP 803.04 and 2434 which indicates that a reasonable number of nucleotide and amino acid sequences are permitted to be claimed in an application.

Applicant would submit that the most recently imposed restriction requirement is decidedly *unreasonable*. Applicant's submit that the sequences in the present application are so interrelated as to be easily searched together without undue burden on the Patent Office. Applicant's have aligned the nucleic acid sequences of the mammary and hippocampal forms of sel-10 so that the examiner may appreciate this fact. In addition the Applicants have aligned SEQ ID NOS:1-10 so that the Examiner may appreciate their considerable degree of identity.

Applicants would respectfully submit that at the very least the previous four way restriction requirement should be reinstated but would also respectfully request that the examiner address the traversal in the previously filed papers.

Respectfully submitted,


Edward F. Rehberg, Attorney
Registration No. 34,703

Date: 8-14-2001

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Section 1

	(1)	1	10	20	30	40	53
hmsel-10	(1)	CTCAGCAGGT	CAGGACAT	TTGGTAGG	GGAAGGTT	GAAAGACAAA	AGCAGCAGG
hhSel-10	(1)	-----	CTCATTAT	TCCCTCGA	GTTC--TT	CTCAGTCAA	--GCTGCATG
Consensus	(1)		C	ATT	T G G	TT	AG CAA GC GCA G

Section 2

	(54)	54	60	70	80	90	106
hmsel-10	(54)	CC	TTGGGT	CTCAGC	CTTTTAAAA	ACTATTAT	TAAATATA
hhSel-10	(40)	TAT--	GTATGT	GTCTCC	GAGAAC	GCGGTT	TGATACTG
Consensus	(54)	T	G	T T G C	AA	TT TA	A T TTT TT

Section 3

	(107)	107	120	130	140	159
hmsel-10	(107)	TAGTGGT	TAGAGC	TTTTAGT	AAATGTGC	CTGTATTAC
hhSel-10	(90)	TACTG--	TGGAG-	TTTTGT	TGCCG	GTTCTGC-TCC
Consensus	(107)	TA TG	T GAG	TTTT T	G CTG	T C T T T G

Section 4

	(160)	160	170	180	190	200	212
hmsel-10	(160)	CAACCA	AGAGGAG	TTTTTAA	AATGTCAA	AACCGGCA	-AAACCTAG
hhSel-10	(139)	CGTG	CCTGAGCA	-TGTCC	CACAT-TAGA	ATCTGTG	ACATACCTAG
Consensus	(160)	C	C	GAG A T T	A AT T	AA C G GA	A ACCTAC CT AA A

Section 5

	(213)	213	220	230	240	250	265
hmsel-10	(212)	TGGCT	TGGTT	CCTGTT	GATCTTAA	AAGTGCA	AAAGAGCC
hhSel-10	(189)	GCTTT	TATATT	GTCTAG	AGACTGCC	AAGCAGC	CGGACAC
Consensus	(213)	G	T TT	GA	AA GC	A A C	CACA AA

Section 6

	(266)	266	280	290	300	318	
hmsel-10	(265)	CCGTGAT	GAAGAT	ATTTAG	CATTAG	CATTCGCC	CAAGGC--CT
hhSel-10	(241)	CAC	TGAAG	GGGAAAA	ATACAG	AAATATG	GGGTTTCT
Consensus	(266)	C	TGA G	GA A TA	A AT	T C A	GGC T T

Section 7

	(319)	319	330	340	350	360	371
hmsel-10	(316)	GTCGA	AGACGGA	-TGAAA	AGAAAG	TTCGAC	CATGGTTCT
hhSel-10	(294)	ATTTTT	TACAAA	ATGAAA	AGAAAG	TTCGAC	CATGGTTCT
Consensus	(319)	T	AC A	TGAAA	AGAAAG	TTCGAC	CATGGTTCT

Section 8

	(372)	372	380	390	400	410	424
hmsel-10	(368)	TTCTTT	TGGGAA	AGAAAC	CATGCAA	AGTCTC	AGAATATA
hhSel-10	(347)	TTCTTT	TGGGAA	AGAAAC	CATGCAA	AGTCTC	AGAATATA
Consensus	(372)	TTCTTT	TGGGAA	AGAAAC	CATGCAA	AGTCTC	AGAATATA

Untitled

		Section 9					
	(425)	425	430	440	450	460	477
hmSel-10	(421)	TTGTACCATGTTTCAGCAACACCAACAACCTTTTGGGGACCTCAGAGCAGCCAAT					
hhSel-10	(400)	TTGTACCATGTTTCAGCAACACCAACAACCTTTTGGGGACCTCAGAGCAGCCAAT					
Consensus	(425)	TTGTACCATGTTTCAGCAACACCAACAACCTTTTGGGGACCTCAGAGCAGCCAAT					
		Section 10					
	(478)	478	490	500	510	520	530
hmSel-10	(474)	GGCCAAGGGCAACAACGACGCCGAATTACATCTGTCCAGGCACCTACAGGCCT					
hhSel-10	(453)	GGCCAAGGGCAACAACGACGCCGAATTACATCTGTCCAGGCACCTACAGGCCT					
Consensus	(478)	GGCCAAGGGCAACAACGACGCCGAATTACATCTGTCCAGGCACCTACAGGCCT					
		Section 11					
	(531)	531	540	550	560	570	583
hmSel-10	(527)	CCAGGAATGGCTAAAAATGTTTCAGAGCTGGAGTGGACCAGAGAAATTGCTTG					
hhSel-10	(506)	CCAGGAATGGCTAAAAATGTTTCAGAGCTGGAGTGGACCAGAGAAATTGCTTG					
Consensus	(531)	CCAGGAATGGCTAAAAATGTTTCAGAGCTGGAGTGGACCAGAGAAATTGCTTG					
		Section 12					
	(584)	584	590	600	610	620	636
hmSel-10	(580)	CTTTAGATGAACTCATTGATAGTTGTGAACCAACACAAGTAAACATATGATG					
hhSel-10	(559)	CTTTAGATGAACTCATTGATAGTTGTGAACCAACACAAGTAAACATATGATG					
Consensus	(584)	CTTTAGATGAACTCATTGATAGTTGTGAACCAACACAAGTAAACATATGATG					
		Section 13					
	(637)	637	650	660	670		689
hmSel-10	(633)	CAAGTGATAGAACCCAGTTTCAACGAGACTTCATTTTCATTGCTCCCTAAAGA					
hhSel-10	(612)	CAAGTGATAGAACCCAGTTTCAACGAGACTTCATTTTCATTGCTCCCTAAAGA					
Consensus	(637)	CAAGTGATAGAACCCAGTTTCAACGAGACTTCATTTTCATTGCTCCCTAAAGA					
		Section 14					
	(690)	690	700	710	720	730	742
hmSel-10	(686)	GTTGGCACTCTATGTGCTTTTCATTCCTGGAACCCAAAGACCTGCTACAAGCAG					
hhSel-10	(665)	GTTGGCACTCTATGTGCTTTTCATTCCTGGAACCCAAAGACCTGCTACAAGCAG					
Consensus	(690)	GTTGGCACTCTATGTGCTTTTCATTCCTGGAACCCAAAGACCTGCTACAAGCAG					
		Section 15					
	(743)	743	750	760	770	780	795
hmSel-10	(739)	CTCAGACATGTCGCTACTGGAGAATTTTGGCTGAAGACAACCTTCTCTGGAGA					
hhSel-10	(718)	CTCAGACATGTCGCTACTGGAGAATTTTGGCTGAAGACAACCTTCTCTGGAGA					
Consensus	(743)	CTCAGACATGTCGCTACTGGAGAATTTTGGCTGAAGACAACCTTCTCTGGAGA					
		Section 16					
	(796)	796	810	820	830		848
hmSel-10	(792)	GAGAAATGCAAAGAAGAGGGGATTGATGAACCATTCACATCAAGAGAAGAAA					
hhSel-10	(771)	GAGAAATGCAAAGAAGAGGGGATTGATGAACCATTCACATCAAGAGAAGAAA					
Consensus	(796)	GAGAAATGCAAAGAAGAGGGGATTGATGAACCATTCACATCAAGAGAAGAAA					

Untitled

							Section 17
	(849)	849	860	870	880	890	901
hmSel-10	(845)	AGTAATAAAAGCAGGTTTCATACACAGTCCATGGAAAAGTGCATACATCAGAC					
hhSel-10	(824)	AGTAATAAAACCAGGTTTCATACACAGTCCATGGAAAAGTGCATACATCAGAC					
Consensus	(849)	AGTAATAAAACCAGGTTTCATACACAGTCCATGGAAAAGTGCATACATCAGAC					
							Section 18
	(902)	902	910	920	930	940	954
hmSel-10	(898)	AGCACAGAATTGATACTAACTGGAGGCGAGGAGAACTCAAATCTCCTAAGGTG					
hhSel-10	(877)	AGCACAGAATTGATACTAACTGGAGGCGAGGAGAACTCAAATCTCCTAAGGTG					
Consensus	(902)	AGCACAGAATTGATACTAACTGGAGGCGAGGAGAACTCAAATCTCCTAAGGTG					
							Section 19
	(955)	955	960	970	980	990	1007
hmSel-10	(951)	CTGAAAGGACATGATGATCATGTGATCACATGCTTACAGTTTTGTGGTAACCG					
hhSel-10	(930)	CTGAAAGGACATGATGATCATGTGATCACATGCTTACAGTTTTGTGGTAACCG					
Consensus	(955)	CTGAAAGGACATGATGATCATGTGATCACATGCTTACAGTTTTGTGGTAACCG					
							Section 20
	(1008)	1008	1020	1030	1040	1050	1060
hmSel-10	(1004)	AATAGTTAGTGGTTCTGATGACAACACTTTAAAAGTTTGGTCAGCAGTCACAG					
hhSel-10	(983)	AATAGTTAGTGGTTCTGATGACAACACTTTAAAAGTTTGGTCAGCAGTCACAG					
Consensus	(1008)	AATAGTTAGTGGTTCTGATGACAACACTTTAAAAGTTTGGTCAGCAGTCACAG					
							Section 21
	(1061)	1061	1070	1080	1090	1100	1113
hmSel-10	(1057)	GCAAATGTCTGAGAACATTAGTGGGACATACAGGTGGAGTATGGTCATCACAA					
hhSel-10	(1036)	GCAAATGTCTGAGAACATTAGTGGGACATACAGGTGGAGTATGGTCATCACAA					
Consensus	(1061)	GCAAATGTCTGAGAACATTAGTGGGACATACAGGTGGAGTATGGTCATCACAA					
							Section 22
	(1114)	1114	1120	1130	1140	1150	1166
hmSel-10	(1110)	ATGAGAGACAACATCATCATTAGTGGATCTACAGATCGGACACTCAAAGTGTG					
hhSel-10	(1089)	ATGAGAGACAACATCATCATTAGTGGATCTACAGATCGGACACTCAAAGTGTG					
Consensus	(1114)	ATGAGAGACAACATCATCATTAGTGGATCTACAGATCGGACACTCAAAGTGTG					
							Section 23
	(1167)	1167	1180	1190	1200		1219
hmSel-10	(1163)	GAATGCAGAGACTGGAGAATGTATACACACCTTATATGGGCATACTTCCACTG					
hhSel-10	(1142)	GAATGCAGAGACTGGAGAATGTATACACACCTTATATGGGCATACTTCCACTG					
Consensus	(1167)	GAATGCAGAGACTGGAGAATGTATACACACCTTATATGGGCATACTTCCACTG					
							Section 24
	(1220)	1220	1230	1240	1250	1260	1272
hmSel-10	(1216)	TGGGTTGTATGCATCTTCATGAAAAAAGAGTTGTTAGCGGTTCTCGAGATGCC					
hhSel-10	(1195)	TGGGTTGTATGCATCTTCATGAAAAAAGAGTTGTTAGCGGTTCTCGAGATGCC					
Consensus	(1220)	TGGGTTGTATGCATCTTCATGAAAAAAGAGTTGTTAGCGGTTCTCGAGATGCC					

						Section 25
(1273)	1273	1280	1290	1300	1310	1325
hmsel-10 (1269)	ACTCTTAGGGTTTGGGATATTGAGACAGGCCAGTGTTTACATGTTTTGATGGG					
hhSel-10 (1248)	ACTCTTAGGGTTTGGGATATTGAGACAGGCCAGTGTTTACATGTTTTGATGGG					
Consensus (1273)	ACTCTTAGGGTTTGGGATATTGAGACAGGCCAGTGTTTACATGTTTTGATGGG					
						Section 26
(1326)	1326	1340	1350	1360		1378
hmsel-10 (1322)	TCATGTTGCAGCAGTCCGCTGTGTTCAATATGATGGCAGGAGGGTTGTTAGTG					
hhSel-10 (1301)	TCATGTTGCAGCAGTCCGCTGTGTTCAATATGATGGCAGGAGGGTTGTTAGTG					
Consensus (1326)	TCATGTTGCAGCAGTCCGCTGTGTTCAATATGATGGCAGGAGGGTTGTTAGTG					
						Section 27
(1379)	1379	1390	1400	1410	1420	1431
hmsel-10 (1375)	GAGCATATGATTTTATGGTAAAGGTGTGGGATCCAGAGACTGAAACCTGTCTA					
hhSel-10 (1354)	GAGCATATGATTTTATGGTAAAGGTGTGGGATCCAGAGACTGAAACCTGTCTA					
Consensus (1379)	GAGCATATGATTTTATGGTAAAGGTGTGGGATCCAGAGACTGAAACCTGTCTA					
						Section 28
(1432)	1432	1440	1450	1460	1470	1484
hmsel-10 (1428)	CACACGTTGCAGGGGCATACTAATAGAGTCTATTTCATTACAGTTTGATGGTAT					
hhSel-10 (1407)	CACACGTTGCAGGGGCATACTAATAGAGTCTATTTCATTACAGTTTGATGGTAT					
Consensus (1432)	CACACGTTGCAGGGGCATACTAATAGAGTCTATTTCATTACAGTTTGATGGTAT					
						Section 29
(1485)	1485	1490	1500	1510	1520	1537
hmsel-10 (1481)	CCATGTGGTGAGTGGATCTCTTGATACATCAATCCGTGTTTGGGATGTGGAGA					
hhSel-10 (1460)	CCATGTGGTGAGTGGATCTCTTGATACATCAATCCGTGTTTGGGATGTGGAGA					
Consensus (1485)	CCATGTGGTGAGTGGATCTCTTGATACATCAATCCGTGTTTGGGATGTGGAGA					
						Section 30
(1538)	1538	1550	1560	1570	1580	1590
hmsel-10 (1534)	CAGGGAATTGCATTACACAGTTAACAGGGCACCAGTCGTTAACAAAGTGGGAATG					
hhSel-10 (1513)	CAGGGAATTGCATTACACAGTTAACAGGGCACCAGTCGTTAACAAAGTGGGAATG					
Consensus (1538)	CAGGGAATTGCATTACACAGTTAACAGGGCACCAGTCGTTAACAAAGTGGGAATG					
						Section 31
(1591)	1591	1600	1610	1620	1630	1643
hmsel-10 (1587)	GAACTCAAAGACAATATTCTTGTCTCTGGGAATGCAGATTCTACAGTTAAAAT					
hhSel-10 (1566)	GAACTCAAAGACAATATTCTTGTCTCTGGGAATGCAGATTCTACAGTTAAAAT					
Consensus (1591)	GAACTCAAAGACAATATTCTTGTCTCTGGGAATGCAGATTCTACAGTTAAAAT					
						Section 32
(1644)	1644	1650	1660	1670	1680	1696
hmsel-10 (1640)	CTGGGATATCAAAACAGGACAGTGTTTACAAACATTGCAAGGTCCCAACAAGC					
hhSel-10 (1619)	CTGGGATATCAAAACAGGACAGTGTTTACAAACATTGCAAGGTCCCAACAAGC					
Consensus (1644)	CTGGGATATCAAAACAGGACAGTGTTTACAAACATTGCAAGGTCCCAACAAGC					

Section 33

	(1697)	1697	1710	1720	1730	1749
hmsel-10 (1693)		ATCAGAGTGCTGTGACCTGTTTACAGTTCAACAAGAAGCTTTGTAATTACCAGC				
hhSel-10 (1672)		ATCAGAGTGCTGTGACCTGTTTACAGTTCAACAAGAAGCTTTGTAATTACCAGC				
Consensus (1697)		ATCAGAGTGCTGTGACCTGTTTACAGTTCAACAAGAAGCTTTGTAATTACCAGC				

Section 34

	(1750)	1750	1760	1770	1780	1790	1802
hmsel-10 (1746)		TCAGATGATGGAAGCTGTAAAGCTATGGGACTTGAAAACGGGTGAATTTATTTCG					
hhSel-10 (1725)		TCAGATGATGGAAGCTGTAAAGCTATGGGACTTGAAAACGGGTGAATTTATTTCG					
Consensus (1750)		TCAGATGATGGAAGCTGTAAAGCTATGGGACTTGAAAACGGGTGAATTTATTTCG					

Section 35

	(1803)	1803	1810	1820	1830	1840	1855
hmsel-10 (1799)		AAACCTAGTCACATTGGAGAGTGGGGGAGTGGGGGAGTTGTGTGGCGGATCA					
hhSel-10 (1778)		AAACCTAGTCACATTGGAGAGTGGGGGAGTGGGGGAGTTGTGTGGCGGATCA					
Consensus (1803)		AAACCTAGTCACATTGGAGAGTGGGGGAGTGGGGGAGTTGTGTGGCGGATCA					

Section 36

	(1856)	1856	1870	1880	1890	1908
hmsel-10 (1852)		GAGCCTCAAACACAAAGCTGGTGTGTGCAGTTGGGAGTCGGAATGGGACTGAA				
hhSel-10 (1831)		GAGCCTCAAACACAAAGCTGGTGTGTGCAGTTGGGAGTCGGAATGGGACTGAA				
Consensus (1856)		GAGCCTCAAACACAAAGCTGGTGTGTGCAGTTGGGAGTCGGAATGGGACTGAA				

Section 37

	(1909)	1909	1920	1930	1940	1950	1961
hmsel-10 (1905)		GAAACCAAGCTGCTGGTGTGCTGGACTTTGATGTGGACATGAAGTGAAGAGCAGA					
hhSel-10 (1884)		GAAACCAAGCTGCTGGTGTGCTGGACTTTGATGTGGACATGAAGTGAAGAGCAGA					
Consensus (1909)		GAAACCAAGCTGCTGGTGTGCTGGACTTTGATGTGGACATGAAGTGAAGAGCAGA					

Section 38

	(1962)	1962	1970	1980	1990	2000	2014
hmsel-10 (1958)		AAAGATGAATTTGTCCAATTGTGTAGACGATATACTCCCTGCCCTTCCCCCTG					
hhSel-10 (1937)		AAAGATGAATTTGTCCAATTGTGTAGACGATATACTCCCTGCCCTTCCCCCTG					
Consensus (1962)		AAAGATGAATTTGTCCAATTGTGTAGACGATATACTCCCTGCCCTTCCCCCTG					

Section 39

	(2015)	2015	2020	2030	2040	2050	2067
hmsel-10 (2011)		CAAAAAGAAAAAAGAAAAAGAAAAAGAAAAAATCCCTTGTTCCTCAGTGGTGC					
hhSel-10 (1990)		CAAAAAGAAAAAAGAAAAAGAAAAAGAAAAAATCCCTTGTTCCTCAGTGGTGC					
Consensus (2015)		CAAAAAGAAAAAAGAAAAAGAAAAAGAAAAAATCCCTTGTTCCTCAGTGGTGC					

Section 40

	(2068)	2068	2080	2090	2100	2110	2120
hmsel-10 (2064)		AGGATGTTGGCTTGGGGCAACAGATTGAAAAGACCTACAGACTAAGAAGGAAA					
hhSel-10 (2043)		AGGATGTTGGCTTGGGGCAACAGATTGAAAAGACCTACAGACTAAGAAGGAAA					
Consensus (2068)		AGGATGTTGGCTTGGGGCAACAGATTGAAAAGACCTACAGACTAAGAAGGAAA					

Untitled

						Section 41
(2121)	2121	2130	2140	2150	2160	2173
hmsel-10 (2117)	AGAAGAAGAGATGACAAACCATAACTGACAAGAGAGGGCGTCTGCTGTCTCATC					
hhSel-10 (2096)	AGAAGAAGAGATGACAAACCATAACTGACAAGAGAGGGCGTCTGCTGTCTCATC					
Consensus (2121)	AGAAGAAGAGATGACAAACCATAACTGACAAGAGAGGGCGTCTGCTGTCTCATC					
						Section 42
(2174)	2174	2180	2190	2200	2210	2226
hmsel-10 (2170)	ACATAAAAGGCTTCACCTTTTGAAGTGAAGGGCAGCTTTGCAAAATGAGACTTTCT					
hhSel-10 (2149)	ACATAAAAGGCTTCACCTTTTGAAGTGAAGGGCAGCTTTGCAAAATGAGACTTTCT					
Consensus (2174)	ACATAAAAGGCTTCACCTTTTGAAGTGAAGGGCAGCTTTGCAAAATGAGACTTTCT					
						Section 43
(2227)	2227	2240	2250	2260		2279
hmsel-10 (2223)	AAATCAAACCAGGTGCAATTATTTCTTTATTTTCTTCTCCAGTGGTCATTGGG					
hhSel-10 (2202)	AAATCAAACCAGGTGCAATTATTTCTTTATTTTCTTCTCCAGTGGTCATTGGG					
Consensus (2227)	AAATCAAACCAGGTGCAATTATTTCTTTATTTTCTTCTCCAGTGGTCATTGGG					
						Section 44
(2280)	2280	2290	2300	2310	2320	2332
hmsel-10 (2276)	GCAGTGTTAATGCTGAAACATCATTACAGATTCTGCTAGCCTGTTCTTTTACC					
hhSel-10 (2255)	GCAGTGTTAATGCTGAAACATCATTACAGATTCTGCTAGCCTGTTCTTTTACC					
Consensus (2280)	GCAGTGTTAATGCTGAAACATCATTACAGATTCTGCTAGCCTGTTCTTTTACC					
						Section 45
(2333)	2333	2340	2350	2360	2370	2385
hmsel-10 (2329)	ACTGACAGCTAGACACCTAGAAAGGAAGTGAATAATATCAAAACAAGTACTG					
hhSel-10 (2308)	ACTGACAGCTAGACACCTAGAAAGGAAGTGAATAATATCAAAACAAGTACTG					
Consensus (2333)	ACTGACAGCTAGACACCTAGAAAGGAAGTGAATAATATCAAAACAAGTACTG					
						Section 46
(2386)	2386	2400	2410	2420		2438
hmsel-10 (2382)	GTTGACTTTCTAATTAGAGAGCATCTGCAACAAAAAGTCATTTTTCTGGAGTG					
hhSel-10 (2361)	GTTGACTTTCTAATTAGAGAGCATCTGCAACAAAAAGTCATTTTTCTGGAGTG					
Consensus (2386)	GTTGACTTTCTAATTAGAGAGCATCTGCAACAAAAAGTCATTTTTCTGGAGTG					
						Section 47
(2439)	2439	2450	2460	2470	2480	2491
hmsel-10 (2435)	GAAAAGCTTAAAAAATTACTGTGAATTGTTTTGTACAGTTATCATGAAAAG					
hhSel-10 (2414)	GAAAAGCTTAAAAAATTACTGTGAATTGTTTTGTACAGTTATCATGAAAAG					
Consensus (2439)	GAAAAGCTTAAAAAATTACTGTGAATTGTTTTGTACAGTTATCATGAAAAG					
						Section 48
(2492)	2492	2500	2510	2520	2530	2544
hmsel-10 (2488)	CTTTTTTTTTTATTTTTTNGCCAACCATTGCCAATGTCAATCAATCACAGTAT					
hhSel-10 (2467)	CTTTTTTTTTTATTTTTTNGCCAACCATTGCCAATGTCAATCAATCACAGTAT					
Consensus (2492)	CTTTTTTTTTTATTTTTTNGCCAACCATTGCCAATGTCAATCAATCACAGTAT					

Untitled

						Section 49
(2545)	2545	2550	2560	2570	2580	2597
hmsel-10 (2541)	TAGCCTCTGTTAATCTATTTACTGTTGCTTCCATATACATTCTTCAATGCATA					
hhSel-10 (2520)	TAGCCTCTGTTAATCTATTTACTGTTGCTTCCATATACATTCTTCAATGCATA					
Consensus (2545)	TAGCCTCTGTTAATCTATTTACTGTTGCTTCCATATACATTCTTCAATGCATA					
						Section 50
(2598)	2598	2610	2620	2630	2640	2650
hmsel-10 (2594)	TGTTGCTCAAAGGTGGCAAGTTGTCCTGGGTTCTGTGAGTCCTGAGATGGATT					
hhSel-10 (2573)	TGTTGCTCAAAGGTGGCAAGTTGTCCTGGGTTCTGTGAGTCCTGAGATGGATT					
Consensus (2598)	TGTTGCTCAAAGGTGGCAAGTTGTCCTGGGTTCTGTGAGTCCTGAGATGGATT					
						Section 51
(2651)	2651	2660	2670	2680	2690	2703
hmsel-10 (2647)	TAATTCTTGATGCTGGTGCTAGAAGTAGGTCTTCAAATATGGGATTGTTGTCC					
hhSel-10 (2626)	TAATTCTTGATGCTGGTGCTAGAAGTAGGTCTTCAAATATGGGATTGTTGTCC					
Consensus (2651)	TAATTCTTGATGCTGGTGCTAGAAGTAGGTCTTCAAATATGGGATTGTTGTCC					
						Section 52
(2704)	2704	2710	2720	2730	2740	2756
hmsel-10 (2700)	CAACCCTGTACTGTACTCCCAGTGGCCAAACTTATTTATGCTGCTAAATGAAA					
hhSel-10 (2679)	CAACCCTGTACTGTACTCCCAGTGGCCAAACTTATTTATGCTGCTAAATGAAA					
Consensus (2704)	CAACCCTGTACTGTACTCCCAGTGGCCAAACTTATTTATGCTGCTAAATGAAA					
						Section 53
(2757)	2757	2770	2780	2790		2809
hmsel-10 (2753)	GAAAGAAAAAGCAAATTATTTTATTTTCTGCTGTGACGTTTATAG					
hhSel-10 (2732)	GAAAGAAAAAGCAAATTATTTTATTTTCTGCTGTGACGTTTATAG					
Consensus (2757)	GAAAGAAAAAGCAAATTATTTTATTTTCTGCTGTGACGTTTATAG					
						Section 54
(2810)	2810	2820	2830	2840	2850	2862
hmsel-10 (2806)	TCCCAGACTGAATTCCAAATTTGCTCTAGTTTGGTTATGGAAAAAGACTTTT					
hhSel-10 (2785)	TCCCAGACTGAATTCCAAATTTGCTCTAGTTTGGTTATGGAAAAAGACTTTT					
Consensus (2810)	TCCCAGACTGAATTCCAAATTTGCTCTAGTTTGGTTATGGAAAAAGACTTTT					
						Section 55
(2863)	2863	2870	2880	2890	2900	2915
hmsel-10 (2859)	TGCCACTGAAACTTGAGCCATCTGTGCCTCTAAGAGGCTGAGAATGGAAGAGT					
hhSel-10 (2838)	TGCCACTGAAACTTGAGCCATCTGTGCCTCTAAGAGGCTGAGAATGGAAGAGT					
Consensus (2863)	TGCCACTGAAACTTGAGCCATCTGTGCCTCTAAGAGGCTGAGAATGGAAGAGT					
						Section 56
(2916)	2916	2930	2940	2950		2968
hmsel-10 (2912)	TTCAGATAATAAAGAGTGAAGTTTGCTGCAAGTAAAGAATTGAGAGTGTGTG					
hhSel-10 (2891)	TTCAGATAATAAAGAGTGAAGTTTGCTGCAAGTAAAGAATTGAGAGTGTGTG					
Consensus (2916)	TTCAGATAATAAAGAGTGAAGTTTGCTGCAAGTAAAGAATTGAGAGTGTGTG					

Untitled

							Section 57
(2969)	2969	2980	2990	3000	3010	3021	
hmsel-10 (2965)	CAAAGCTTATTTTCTTTTATCTGGGC AAAAATTAAAACACATTCCTTGGGAACA						
hhSel-10 (2944)	CAAAGCTTATTTTCTTTTATCTGGGC AAAAATTAAAACACATTCCTTGGGAACA						
Consensus (2969)	CAAAGCTTATTTTCTTTTATCTGGGC AAAAATTAAAACACATTCCTTGGGAACA						
							Section 58
(3022)	3022	3030	3040	3050	3060	3074	
hmsel-10 (3018)	GAGCTATTACTTGCCCTGTTCTGTGGAGAACTTTTCTTTTGGAGGGCTGTGGT						
hhSel-10 (2997)	GAGCTATTACTTGCCCTGTTCTGTGGAGAACTTTTCTTTTGGAGGGCTGTGGT						
Consensus (3022)	GAGCTATTACTTGCCCTGTTCTGTGGAGAACTTTTCTTTTGGAGGGCTGTGGT						
							Section 59
(3075)	3075	3080	3090	3100	3110	3127	
hmsel-10 (3071)	GAATGGATGAACGTACATCGTAAAACTGACAAAATATTTTAAAAATATATAAA						
hhSel-10 (3050)	GAATGGATGAACGTACATCGTAAAACTGACAAAATATTTTAAAAATATATAAA						
Consensus (3075)	GAATGGATGAACGTACATCGTAAAACTGACAAAATATTTTAAAAATATATAAA						
							Section 60
(3128)	3128	3140	3150	3160	3170	3180	
hmsel-10 (3124)	ACACAAAATTAAAAATAAAGTTGCTGGTCAGTCTTAGTGTTTTACAGTATTTGG						
hhSel-10 (3103)	ACACAAAATTAAAAATAAAGTTGCTGGTCAGTCTTAGTGTTTTACAGTATTTGG						
Consensus (3128)	ACACAAAATTAAAAATAAAGTTGCTGGTCAGTCTTAGTGTTTTACAGTATTTGG						
							Section 61
(3181)	3181	3190	3200	3210	3220	3233	
hmsel-10 (3177)	GAAAACAAC TGTTACAGT TTTAT TGCTCTGAGTAACTGACAAAGCAGAAACTA						
hhSel-10 (3156)	GAAAACAAC TGTTACAGT TTTAT TGCTCTGAGTAACTGACAAAGCAGAAACTA						
Consensus (3181)	GAAAACAAC TGTTACAGT TTTAT TGCTCTGAGTAACTGACAAAGCAGAAACTA						
							Section 62
(3234)	3234	3240	3250	3260	3270	3286	
hmsel-10 (3230)	TTCAGTTTTTG TAGTAAAGGCGTCACATGCAAACAAACAAAATGAATGAAACA						
hhSel-10 (3209)	TTCAGTTTTTG TAGTAAAGGCGTCACATGCAAACAAACAAAATGAATGAAACA						
Consensus (3234)	TTCAGTTTTTG TAGTAAAGGCGTCACATGCAAACAAACAAAATGAATGAAACA						
							Section 63
(3287)	3287	3300	3310	3320	3339		
hmsel-10 (3283)	GTCAAATGGTTTGCCCTCATTCTCCAAGAGCCACAAC TCAAGCTGAAC TGTGAA						
hhSel-10 (3262)	GTCAAATGGTTTGCCCTCATTCTCCAAGAGCCACAAC TCAAGCTGAAC TGTGAA						
Consensus (3287)	GTCAAATGGTTTGCCCTCATTCTCCAAGAGCCACAAC TCAAGCTGAAC TGTGAA						
							Section 64
(3340)	3340	3350	3360	3370	3380	3392	
hmsel-10 (3336)	AGTGGTTTAAACACTGTATCCTAGGCGATCTTTTTTCTCCTTCTGTTTATTTT						
hhSel-10 (3315)	AGTGGTTTAAACACTGTATCCTAGGCGATCTTTTTTCTCCTTCTGTTTATTTT						
Consensus (3340)	AGTGGTTTAAACACTGTATCCTAGGCGATCTTTTTTCTCCTTCTGTTTATTTT						

Untitled

Section 65

	(3393)	3393	3400	3410	3420	3430	3445
hmsel-10	(3389)	TTTGNTTGTTTTATTTATAGTCTGATTAAACAATCAGATTCAAGTTGGTTA					
hhSel-10	(3368)	TTTGNTTGTTTTATTTATAGTCTGATTAAACAATCAGATTCAAGTTGGTTA					
Consensus	(3393)	TTTGNTTGTTTTATTTATAGTCTGATTAAACAATCAGATTCAAGTTGGTTA					

Section 66

	(3446)	3446	3460	3470	3480	3498
hmsel-10	(3442)	ATTTTAGTTATGTAACAACCTGACATGATGGAGGAAAACAACCTTTAAAGGGA				
hhSel-10	(3421)	ATTTTAGTTATGTAACAACCTGACATGATGGAGGAAAACAACCTTTAAAGGGA				
Consensus	(3446)	ATTTTAGTTATGTAACAACCTGACATGATGGAGGAAAACAACCTTTAAAGGGA				

Section 67

	(3499)	3499	3510	3520	3530	3540	3551
hmsel-10	(3495)	TTGTGTCTATGGTTTGATTCACCTTAGAAATTTATTTTCTTATAACTTAAGTG					
hhSel-10	(3474)	TTGTGTCTATGGTTTGATTCACCTTAGAAATTTATTTTCTTATAACTTAAGTG					
Consensus	(3499)	TTGTGTCTATGGTTTGATTCACCTTAGAAATTTATTTTCTTATAACTTAAGTG					

Section 68

	(3552)	3552	3560	3575
hmsel-10	(3548)	CAATAAAATGTGTTTTTTCATGTT		
hhSel-10	(3527)	CAATAAAATGTGTTTTTTCATGTT		
Consensus	(3552)	CAATAAAATGTGTTTTTTCATGTT		

Section 1

	(1)	1	10	20	30	40	53
hhSel-10-(1)	3	(1)	MCVPRSGILILSCICLYCGVLLPVLLPNLPFLTCLSMSTLESVTYLPEKGLYCQ				
hhSel-10-(2)	4	(1)	-----MSTLESVTYLPEKGLYCQ				
hhSel-10-(3)	5	(1)	-----				
hhSel-10-(4)	6	(1)	-----				
hhSel-10-(5)	7	(1)	-----				
hmsel-10-(1)	8	(1)	-----MSKPGKPTLNHGLVP				
hmsel-10-(2)	9	(1)	-----				
hmsel-10-(3)	10	(1)	-----				
Consensus	(1)						

Section 2

	(54)	54	60	70	80	90	106
hhSel-10-(1)	(54)	RLPSSRTHGGTESLKGKNTENMGFYGT	LKMTFYKMKRKLDHGSEVR	SFSLGKK			
hhSel-10-(2)	(19)	RLPSSRTHGGTESLKGKNTENMGFYGT	LKMTFYKMKRKLDHGSEVR	SFSLGKK			
hhSel-10-(3)	(1)	-----MGFYGT	LKMTFYKMKRKLDHGSEVR	SFSLGKK			
hhSel-10-(4)	(1)	-----M	FYKMKRKLDHGSEVR	SFSLGKK			
hhSel-10-(5)	(1)	-----M	FYKMKRKLDHGSEVR	SFSLGKK			
hmsel-10-(1)	(16)	VDLKSACEPLPHQTVMKIFSISIIAQGLPFCRR	MRKLDHGSEVR	SFSLGKK			
hmsel-10-(2)	(1)	-----MKIFSISIIAQGLPFCRR	MRKLDHGSEVR	SFSLGKK			
hmsel-10-(3)	(1)	-----M	KRKLDHGSEVR	SFSLGKK			
Consensus	(54)		K		L	MIFYKMKRKLDHGSEVR	SFSLGKK

Section 3

	(107)	107	120	130	140	159
hhSel-10-(1)	(107)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
hhSel-10-(2)	(72)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
hhSel-10-(3)	(33)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
hhSel-10-(4)	(25)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
hhSel-10-(5)	(25)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
hmsel-10-(1)	(69)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
hmsel-10-(2)	(39)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
hmsel-10-(3)	(20)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			
Consensus	(107)	PCKVSEYTS	TTGLVPCSATPTTFGDLRAANGQGQORRRITSVQPPTGLQEWLK			

Section 4

	(160)	160	170	180	190	200	212
hhSel-10-(1)	(160)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
hhSel-10-(2)	(125)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
hhSel-10-(3)	(86)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
hhSel-10-(4)	(78)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
hhSel-10-(5)	(78)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
hmsel-10-(1)	(122)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
hmsel-10-(2)	(92)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
hmsel-10-(3)	(73)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					
Consensus	(160)	MFQSWGPEKLLALDELIDSCEPTQVKHMMQVIEPQFORDFISLLPKELALYV					

Section 5

	(213)	213	220	230	240	250	265
hhSel-10-(1)	(213)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
hhSel-10-(2)	(178)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
hhSel-10-(3)	(139)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
hhSel-10-(4)	(131)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
hhSel-10-(5)	(131)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
hmsel-10-(1)	(175)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
hmsel-10-(2)	(145)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
hmsel-10-(3)	(126)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					
Consensus	(213)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVIKPG					

Section 6

	(266)	266	280	290	300	318
hhSel-10-(1)	(266)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
hhSel-10-(2)	(231)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
hhSel-10-(3)	(192)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
hhSel-10-(4)	(184)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
hhSel-10-(5)	(184)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
hmsel-10-(1)	(228)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
hmsel-10-(2)	(198)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
hmsel-10-(3)	(179)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				
Consensus	(266)	FIHSPWKSAYIROHRIDTNWRRGELKSPKVLKGHDDHVTCLQFCGNRIVSGS				

Section 7

	(319)	319	330	340	350	360	371
hhSel-10-(1)	(319)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
hhSel-10-(2)	(284)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
hhSel-10-(3)	(245)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
hhSel-10-(4)	(237)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
hhSel-10-(5)	(237)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
hmsel-10-(1)	(281)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
hmsel-10-(2)	(251)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
hmsel-10-(3)	(232)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					
Consensus	(319)	DDNTLKVWSAVTGKCLRTL VGHTGGVWSSQMRDNIITSGSTDRTLKVWNAETG					

Section 8

	(372)	372	380	390	400	410	424
hhSel-10-(1)	(372)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
hhSel-10-(2)	(337)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
hhSel-10-(3)	(298)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
hhSel-10-(4)	(290)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
hhSel-10-(5)	(290)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
hmsel-10-(1)	(334)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
hmsel-10-(2)	(304)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
hmsel-10-(3)	(285)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					
Consensus	(372)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVL MGHVAAV					

Section 9

	(425)	425	430	440	450	460	477
hhSel-10-(1)	(425)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
hhSel-10-(2)	(390)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
hhSel-10-(3)	(351)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
hhSel-10-(4)	(343)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
hhSel-10-(5)	(343)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
hmsel-10-(1)	(387)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
hmsel-10-(2)	(357)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
hmsel-10-(3)	(338)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ
Consensus	(425)	RCVQYD	GRRVVS	SGAYDF	FMVKVW	DPETET	CLHTLQ

Section 10

	(478)	478	490	500	510	520	530
hhSel-10-(1)	(478)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
hhSel-10-(2)	(443)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
hhSel-10-(3)	(404)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
hhSel-10-(4)	(396)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
hhSel-10-(5)	(396)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
hmsel-10-(1)	(440)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
hmsel-10-(2)	(410)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
hmsel-10-(3)	(391)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV
Consensus	(478)	SLDTSIR	VWDVET	GNCHTL	TGHQSL	TSGMEL	KDNILV

Section 11

	(531)	531	540	550	560	570	583
hhSel-10-(1)	(531)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
hhSel-10-(2)	(496)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
hhSel-10-(3)	(457)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
hhSel-10-(4)	(449)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
hhSel-10-(5)	(449)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
hmsel-10-(1)	(493)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
hmsel-10-(2)	(463)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
hmsel-10-(3)	(444)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK
Consensus	(531)	GQCLQTL	QGPKNH	QSAVTC	LQFNKN	FVITSS	DDGTVK

Section 12

	(584)	584	590	600	610	628
hhSel-10-(1)	(584)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
hhSel-10-(2)	(549)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
hhSel-10-(3)	(510)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
hhSel-10-(4)	(502)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
hhSel-10-(5)	(502)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
hmsel-10-(1)	(546)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
hmsel-10-(2)	(516)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
hmsel-10-(3)	(497)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE
Consensus	(584)	ESGGSGG	VVWRIR	ASNTKL	VCAVGS	SRNGTE